STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:	10/565,/64
Source:	IFUP,
Date Processed by STIC:	1/27/06
	7

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FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

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http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm

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Revised 01/10/06



IFWP

RAW SEQUENCE LISTING

DATE: 01/27/2006

PATENT APPLICATION: US/10/565,164

TIME: 09:22:22

Input Set : A:\Q92717 Sequence Listing.txt Output Set: N:\CRF4\01272006\J565164.raw

5 <110 > APPLICANT: POSTECH FOUNDATION

7 <120> TITLE OF INVENTION: INOSITOL-BASED MOLECULAR TRANSPORTERS AND PROCESSES FOR THE

PREPARATION THEREOF

10 <130> FILE REFERENCE: Q92717

C--> 12 <140> CURRENT APPLICATION NUMBER: US/10/565,164

C--> 12 <141> CURRENT FILING DATE: 2006-01-19

12 <150> PRIOR APPLICATION NUMBER: KR10-2004-0014833

13 <151> PRIOR FILING DATE: 2004-03-05

15 <160> NUMBER OF SEO ID NOS: 8

17 <170> SOFTWARE: KopatentIn 1.71

ERRORED SEQUENCES

Does Not Comply 131 <210> SEQ ID NO: 8 Corrected Diskette Needed 132 <211> LENGTH: 17 location 35 to 49
location 35 to 49
account for
g Gly 15 amino acids.

What about the
remaining 2 amino
acids? (serie
this sequence
for 17 amino
acids) 133 <212> TYPE: PRT 134 <213 > ORGANISM: Artificial Sequence 136 <220> FEATURE: 137 <223 > OTHER INFORMATION, a.a. 35 to 49 of FHV Coat 140 <400> SEQUENCE: 8 141 Arg Arg Arg Asn Arg Thr Arg Arg Asn Arg Arg Arg Val Arg Arg Gly 142 144 Cys E--> 149/

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/565,164

DATE: 01/27/2006 TIME: 09:22:23

Input Set : A:\Q92717 Sequence Listing.txt
Output Set: N:\CRF4\01272006\J565164.raw

L:12 M:270 C: Current Application Number differs, Replaced Current Application No

L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date

L:149 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:8